RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/807,517 A_
Source: _FCW/6_
Date Processed by STIC: _7/13/06_

ENTERED

RAW SEQUENCE LISTING   DATE: 07/13/2006

PATENT APPLICATION: US/10/807,517A   TIME: 08:43:40

Input Set : **F:\P-2762-US3.txt**

Output Set: **N:\CRF4\07132006\J807517A.raw**

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   3 <110> APPLICANT: GTx, Inc.
   4        et al.,, Steiner
   5        Steiner, Et al.,
   7 <120> TITLE OF INVENTION: ISOLATED NUCLEIC ACIDS ENCODING RAT P-HYDE PROTEIN
   9 <130> FILE REFERENCE: P-2762-US3
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/807,517A
C--> 11 <141> CURRENT FILING DATE: 2004-03-24
  11 <160> NUMBER OF SEQ ID NOS: 11
  13 <170> SOFTWARE: PatentIn version 3.3
  15 <210> SEQ ID NO: 1
  16 <211> LENGTH: 1886
  17 <212> TYPE: DNA
  18 <213> ORGANISM: Human
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  25 cccgatgagg cccccaaagt gagcatcctg ggtagcgggg actttgcccg ctccctggcc    180
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  35 catcgtgagt ccaatgctga gtacctggcc tccctcttcc ccacttgcac agtggtcaag    480
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102          35                  40                  45
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106      50                  55                  60
109 Ala Arg Leu Phe Pro Ser Ala Ala Gln Val Thr Phe Gln Glu Glu Ala
110 65                  70                  75                  80
113 Val Ser Ser Pro Glu Val Ile Phe Val Ala Val Phe Arg Glu His Tyr
114                  85                  90                  95
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118              100                 105                 110
121 Asp Val Ser Asn Pro Thr Glu Gln Glu His Leu Gln His Arg Glu Ser
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129 Ala Phe Asn Val Ile Ser Ala Trp Thr Leu Gln Ala Gly Pro Arg Asp
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134                 165                 170                 175
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142          195                 200                 205
145 Leu Pro Ala Trp Lys Val Pro Thr Leu Leu Ala Leu Gly Leu Phe Val
146      210                 215                 220
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150 225                 230                 235                 240
153 Gln Glu Ser Gln Asn Lys Phe Phe Lys Leu Pro Val Ser Val Val Asn
154              245                 250                 255
157 Thr Thr Leu Pro Cys Val Ala Tyr Val Leu Leu Ser Leu Val Tyr Leu
158              260                 265                 270
161 Pro Gly Val Leu Ala Ala Ala Leu Gln Leu Arg Arg Gly Thr Lys Tyr
162      275                 280                 285
165 Gln Arg Phe Pro Asp Trp Leu Asp His Trp Leu Gln His Arg Lys Gln
166      290                 295                 300
169 Ile Gly Leu Leu Ser Phe Phe Cys Ala Ala Leu His Ala Leu Tyr Ser
170 305                 310                 315                 320
173 Phe Cys Leu Pro Leu Arg Arg Ala His Arg Tyr Asp Leu Val Asn Leu
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182             355              360              365
185 Thr Leu Ser Leu Leu Ala Val Thr Ser Leu Pro Ser Ile Ala Asn Ser
186         370              375              380
189 Leu Asn Trp Arg Glu Phe Ser Phe Val Gln Ser Ser Leu Gly Phe Val
190 385                390              395                400
193 Ala Leu Val Leu Ser Thr Leu His Thr Leu Thr Tyr Gly Trp Thr Arg
194             405              410              415
197 Ala Phe Glu Glu Ser Arg Tyr Lys Phe Tyr Leu Pro Pro Thr Phe Thr
198         420              425              430
201 Leu Thr Leu Leu Val Pro Cys Val Val Ile Leu Ala Lys Ala Leu Phe
202         435              440              445
205 Leu Leu Pro Cys Ile Ser Arg Arg Leu Ala Arg Ile Arg Arg Gly Trp
206     450              455              460
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219 <212> TYPE: DNA
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227 cccgatgagg cccccaaagt gagcatcctg ggtagcgggg actttgcccg ctccctggcc   180
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231 gccaggctgt ttccctcagc ggcccaagtg actttccaag aggaggcagt gagctccccg   300
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247 tgcttctatg cctacaactt cgtccgggac gttctgcagc cctatgtgca ggaaagccag   780
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289 gctaaaaagt tgggtctctg agatttcaac ttgtagattt aaaaacaagt gccgtacgtt  2040
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312             35                  40                  45
315 Gly Ser Gly Phe Lys Val Val Val Gly Ser Arg Asn Pro Lys Arg Thr
316         50                  55                  60
319 Ala Arg Leu Phe Pro Ser Ala Ala Gln Val Thr Phe Gln Glu Glu Ala
320 65                  70                  75                  80
323 Val Ser Ser Pro Glu Val Ile Phe Val Ala Val Phe Arg Glu His Tyr
324                 85                  90                  95
327 Ser Ser Leu Cys Ser Leu Ser Asp Gln Leu Ala Gly Lys Ile Leu Val
328             100                 105                 110
331 Asp Val Ser Asn Pro Thr Glu Gln Glu His Leu Gln His Arg Glu Ser
332         115                 120                 125
335 Asn Ala Glu Tyr Leu Ala Ser Leu Phe Pro Thr Cys Thr Val Val Lys
336     130                 135                 140
339 Ala Phe Asn Val Ile Ser Ala Trp Thr Leu Gln Ala Gly Pro Arg Asp
340 145                 150                 155                 160
343 Gly Asn Gly Gln Val Pro Ile Cys Gly Asp Gln Pro Glu Ala Lys Arg
344             165                 170                 175
347 Ala Val Ser Glu Met Ala Leu Ala Met Gly Phe Met Pro Val Asp Met
348             180                 185                 190
351 Gly Ser Leu Ala Ser Ala Trp Glu Val Glu Ala Met Pro Leu Arg Leu
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355 Leu Pro Ala Trp Lys Val Pro Thr Leu Leu Ala Leu Gly Leu Phe Val
356     210                 215                 220
359 Cys Phe Tyr Ala Tyr Asn Phe Val Arg Asp Val Leu Gln Pro Tyr Val
360 225                 230                 235                 240
363 Gln Glu Ser Gln Asn Lys Phe Phe Lys Leu Pro Val Ser Val Val Asn
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364                    245                  250                     255
367 Thr Thr Leu Pro Cys Val Ala Tyr Val Leu Leu Ser Leu Val Tyr Leu
368                    260                  265                 270
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372             275                  280                 285
375 Gln Arg Phe Pro Asp Trp Leu Asp His Trp Leu Gln His Arg Lys Gln
376         290                  295                 300
379 Ile Gly Leu Leu Ser Phe Phe Cys Ala Ala Leu His Ala Leu Tyr Ser
380 305                  310                  315                 320
383 Phe Cys Leu Pro Leu Arg Arg Ala His Arg Tyr Asp Leu Val Asn Leu
384             325                  330                 335
387 Ala Val Lys Gln Val Leu Ala Asn Lys Ser His Leu Trp Val Glu Glu
388             340                  345                 350
391 Val Trp Arg Met Glu Ile Tyr Leu Ser Leu Gly Val Leu Ala Leu Gly
392         355                  360                 365
395 Thr Leu Ser Leu Leu Ala Val Thr Ser Leu Pro Ser Ile Ala Asn Ser
396         370                  375                 380
399 Leu Asn Trp Arg Glu Phe Ser Phe Val Gln Cys Val Ala Thr Ser Ser
400 385                  390                  395                 400
403 Ala Gly Asn Thr Gly Ser Gly Thr Arg Arg Pro Glu Ser Gln Ser Gln
404             405                  410                 415
407 Asp Pro His Leu Pro Ala Pro His His Gln Thr Ser Phe Leu Gly Pro
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 07/13/2006
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Input Set : **F:\P-2762-US3.txt**
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<u>Invalid <213> Response:</u>
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:10,11

VERIFICATION SUMMARY   DATE: 07/13/2006
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Input Set : **F:\P-2762-US3.txt**
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L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date